

10/522366

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SEQUENCE LISTING

<110> National Institute of Advanced Industrial Science and Technology

<120> Lethal gene markers for transformant selection

<130> -23603055

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<160> 24

<170> PatentIn Ver. 2.1

<210> 1

<211> 28

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<400> 1

gctgatgctg cattgaggttc tgcttatgg

28

<210> 2

<211> 57

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<400> 2

gttaaatcca atttaagtcc cataacttgg ccgctatggc ctcaaagata tttcttg 57

<210> 3

<211> 57

<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence:primer

<400> 3

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<210> 4

<211> 28

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<400> 4

tcatccctga taatatttga tcaccaat

28

<210> 5

<211> 43

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<400> 5

gcatggccgc ctggccgaa agttttaaa gattacggc atg

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<210> 6

<211> 34

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<400> 6

cgatgaattc tcaccaatca ccatcacgat aatc

34

<210> 7

<211> 598

<212> DNA

<213> E.coli

<400> 7

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cagaatggtg gtggaaaacg caagcgctgg actggagata aagggcgtaa gatttatgag 180
tgggattctc agcatggtga gcttggggg tatcgtgcc a gtatggtca gcatcttggc 240
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aaatatctt gaggccatag cggccaagtt atggactta aattggattt aacttggttt 360
gataaaagta cagaagattt taagggtgag gagtattcaa aagattttgg agatgacggt 420
tcagttatgg aaagtctagg tgtgccttt aaggataatg ttaataacgg ttgccttgat 480
gttatagctg aatgggtacc tttgctacaa ccataactta atcatcaa at tgatattcc 540
gataatgagt atttgttc gtttgattat cgtgatggtg attggtgaga attcatcg 598

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<211> 40

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<213> Artificial Sequence

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<223> Description of Artificial Sequence:primer

<400> 8

tagtagtagt agtagaaagg ttttaagat tacgggcatg

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<210> 9

<211> 46

<212> DNA

<213> E.coli

<400> 9

gcatggccgc ctggccgta gaaaggttt aaagattacg ggcatt

46

<210> 10

<211> 49

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

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49

<210> 11

<211> 52

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<400> 11

gcatggccgc ctcggccgta gtagtagaaa ggttttaag attacggca tg

52

<210> 12

<211> 55

<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence:primer

<400> 12

gcatggccgc ctcggccgta gtagtagtag aaaggttta aagattacgg gcatg

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<210> 13

<211> 58

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:primer

<400> 13

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<210> 14

<211> 607

<212> DNA

<213> E.coli

<400> 14

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tgctttagt ttatagctga atggtagtacct ttgctacaac catactttaa tcataaaattt 540

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<210> 15

<211> 258

<212> DNA

<213> E.coli

<400> 15

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aaggataatg ttaataacgg ttgccttgat gttatacgctg aatgggtacc ttgcataaa 180
ccatactta atcatcaaat tgatattcc gataatgagt attttgttcc gtttgattat 240
cgtgatggtg attggta 258

<210> 16

<211> 3066

<212> DNA

<213> E.coli

<400> 16

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cccgccaaca catcacgggc cacaaaattt tttgtggccc gctctgcgtt ttctaagtgt 180
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cgatcg 3066

<210> 17

<211> 551

<212> PRT

<213> E.coli

<400> 17

Met Ser Gly Gly Asp Gly Arg Gly His Asn Thr Gly Ala His Ser Thr

1

5

10

15

Ser Gly Asn Ile Asn Gly Gly Pro Thr Gly Leu Gly Val Gly Gly

20

25

30

Ala Ser Asp Gly Ser Gly Trp Ser Ser Glu Asn Asn Pro Trp Gly Gly

35

40

45

Gly Ser Gly Ser Gly Ile His Trp Gly Gly Ser Gly His Gly Asn

50

55

60

Gly Gly Gly Asn Gly Asn Ser Gly Gly Ser Gly Thr Gly Gly Asn

65

70

75

80

Leu Ser Ala Val Ala Ala Pro Val Ala Phe Gly Phe Pro Ala Leu Ser

85

90

95

Thr Pro Gly Ala Gly Gly Leu Ala Val Ser Ile Ser Ala Gly Ala Leu

100

105

110

Ser Ala Ala Ile Ala Asp Ile Met Ala Ala Leu Lys Gly Pro Phe Lys

115

120

125

Phe Gly Leu Trp Gly Val Ala Leu Tyr Gly Val Leu Pro Ser Gln Ile

130 135 140

Ala Lys Asp Asp Pro Asn Met Met Ser Lys Ile Val Thr Ser Leu Pro

145 150 155 160

Ala Asp Asp Ile Thr Glu Ser Pro Val Ser Ser Leu Pro Leu Asp Lys

165 170 175

Ala Thr Val Asn Val Asn Val Arg Val Val Asp Asp Val Lys Asp Glu

180 185 190

Arg Gln Asn Ile Ser Val Val Ser Gly Val Pro Met Ser Val Pro Val

195 200 205

Val Asp Ala Lys Pro Thr Glu Arg Pro Gly Val Phe Thr Ala Ser Ile

210 215 220

Pro Gly Ala Pro Val Leu Asn Ile Ser Val Asn Asn Ser Thr Pro Ala

225 230 235 240

Val Gln Thr Leu Ser Pro Gly Val Thr Asn Asn Thr Asp Lys Asp Val

245 250 255

Arg Pro Ala Gly Phe Thr Gln Gly Gly Asn Thr Arg Asp Ala Val Ile

260 265 270

Arg Phe Pro Lys Asp Ser Gly His Asn Ala Val Tyr Val Ser Val Ser

275

280

285

Asp Val Leu Ser Pro Asp Gln Val Lys Gln Arg Gln Asp Glu Glu Asn

290

295

300

Arg Arg Gln Gln Glu Trp Asp Ala Thr His Pro Val Glu Ala Ala Glu

305

310

315

320

Arg Asn Tyr Glu Arg Ala Arg Ala Glu Leu Asn Gln Ala Asn Glu Asp

325

330

335

Val Ala Arg Asn Gln Glu Arg Gln Ala Lys Ala Val Gln Val Tyr Asn

340

345

350

Ser Arg Lys Ser Glu Leu Asp Ala Ala Asn Lys Thr Leu Ala Asp Ala

355

360

365

Ile Ala Glu Ile Lys Gln Phe Asn Arg Phe Ala His Asp Pro Met Ala

370

375

380

Gly Gly His Arg Met Trp Gln Met Ala Gly Leu Lys Ala Gln Arg Ala

385

390

395

400

Gln Thr Asp Val Asn Asn Lys Gln Ala Ala Phe Asp Ala Ala Lys

405

410

415

Glu Lys Ser Asp Ala Asp Ala Ala Leu Ser Ser Ala Met Glu Ser Arg

420

425

430

Lys Lys Lys Glu Asp Lys Lys Arg Ser Ala Glu Asn Asn Leu Asn Asp

435

440

445

Glu Lys Asn Lys Pro Arg Lys Gly Phe Lys Asp Tyr Gly His Asp Tyr

450

455

460

His Pro Ala Pro Lys Thr Glu Asn Ile Lys Gly Leu Gly Asp Leu Lys

465

470

475

480

Pro Gly Ile Pro Lys Thr Pro Lys Gln Asn Gly Gly Lys Arg Lys

485

490

495

Arg Trp Thr Gly Asp Lys Gly Arg Lys Ile Tyr Glu Trp Asp Ser Gln

500

505

510

His Gly Glu Leu Glu Gly Tyr Arg Ala Ser Asp Gly Gln His Leu Gly

515

520

525

Ser Phe Asp Pro Lys Thr Gly Asn Gln Leu Lys Gly Pro Asp Pro Lys

530

535

540

Arg Asn Ile Lys Lys Tyr Leu

545

550

<210> 18

<211> 110

<212> PRT

<213> E.coli

<400> 18

Ala Glu Asn Asn Leu Asn Asp Glu Lys Asn Lys Pro Arg Lys Gly Phe

1

5

10

15

Lys Asp Tyr Gly His Asp Tyr His Pro Ala Pro Lys Thr Glu Asn Ile

20

25

30

Lys Gly Leu Gly Asp Leu Lys Pro Gly Ile Pro Lys Thr Pro Lys Gln

35

40

45

Asn Gly Gly Lys Arg Lys Arg Trp Thr Gly Asp Lys Gly Arg Lys

50

55

60

Ile Tyr Glu Trp Asp Ser Gln His Gly Glu Leu Glu Gly Tyr Arg Ala

65

70

75

80

Ser Asp Gly Gln His Leu Gly Ser Phe Asp Pro Lys Thr Gly Asn Gln

85

90

95

Leu Lys Gly Pro Asp Pro Lys Arg Asn Ile Lys Lys Tyr Leu

100

105

110

<210> 19

<211> 97

<212> PRT

<213> E.coli

<400> 19

Lys Gly Phe Lys Asp Tyr Gly His Asp Tyr His Pro Ala Pro Lys Thr
1 5 10 15

Glu Asn Ile Lys Gly Leu Gly Asp Leu Lys Pro Gly Ile Pro Lys Thr
20 25 30

Pro Lys Gln Asn Gly Gly Lys Arg Lys Arg Trp Thr Gly Asp Lys
35 40 45

Gly Arg Lys Ile Tyr Glu Trp Asp Ser Gln His Gly Glu Leu Glu Gly
50 55 60

Tyr Arg Ala Ser Asp Gly Gln His Leu Gly Ser Phe Asp Pro Lys Thr
65 70 75 80

Gly Asn Gln Leu Lys Gly Pro Asp Pro Lys Arg Asn Ile Lys Lys Tyr
85 90 95

Leu

<210> 20

<211> 330

<212> DNA

<213> E.coli

<400> 1

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caaaggcagaa tggtggtgga aaacgcaagc gctggactgg agataaagg cgtaagattt 180
atgagtggaa ttctcagcat ggtgagcttg aggggtatcg tgccagtatcg ggtcagcatc 240
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tcaagaaata tctttgaggc catagcggcc 330

<210> 21

<211> 60

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:adapter

<400> 2

gatccccggg taccgaggcc gcctcgcccg agctcgaatt cggccggcca tagcggccgc 60

<210> 22

<211> 60

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:adapter

<400> 3

aattgcggcc gctatggccg gccgaattcg agctcggccg aggccgcctc ggtacccggg 60

<210> 23

<211> 650

<212> DNA

<213> S.cerevisiae

<400> 4

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gctactctcc caaaaccaaa aggtctccgc tgacttaggc acatctgaca gaagtggaaat 180
caaggctaga aagactggaa cagctatttc tactgatttt tcctcgagaa gacctgaca 240
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acatgtcgga tggcttgcgc ttcctgaaaa cggaccccaa caataatggg ttctttggcg 600
acggttctct cttatgtatt ctgcgtgac tgactgaggc catagcggcc 650

<210> 24

<211> 535

<212> DNA

<213> A. oryzae

<400> 5

ggccgcctcg gccattacta gtctactagt aactctgtct tatcgtcatc tcccataggt 60
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